

FIG.1

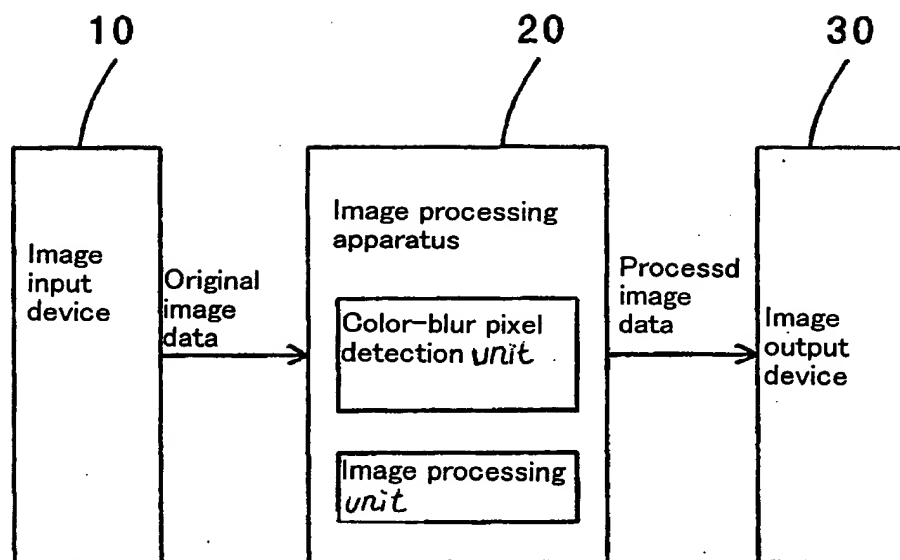


FIG. 2

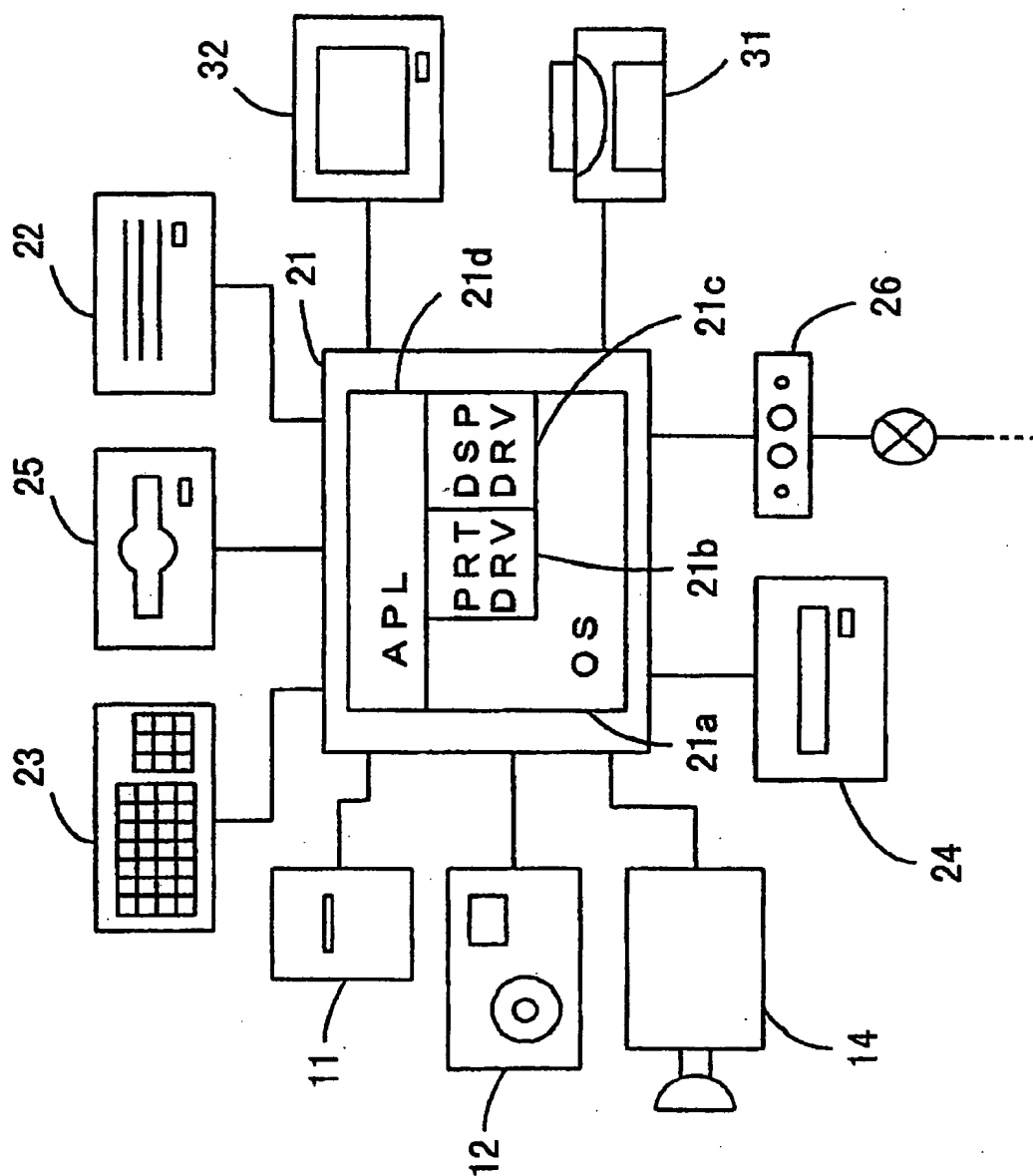


FIG. 3

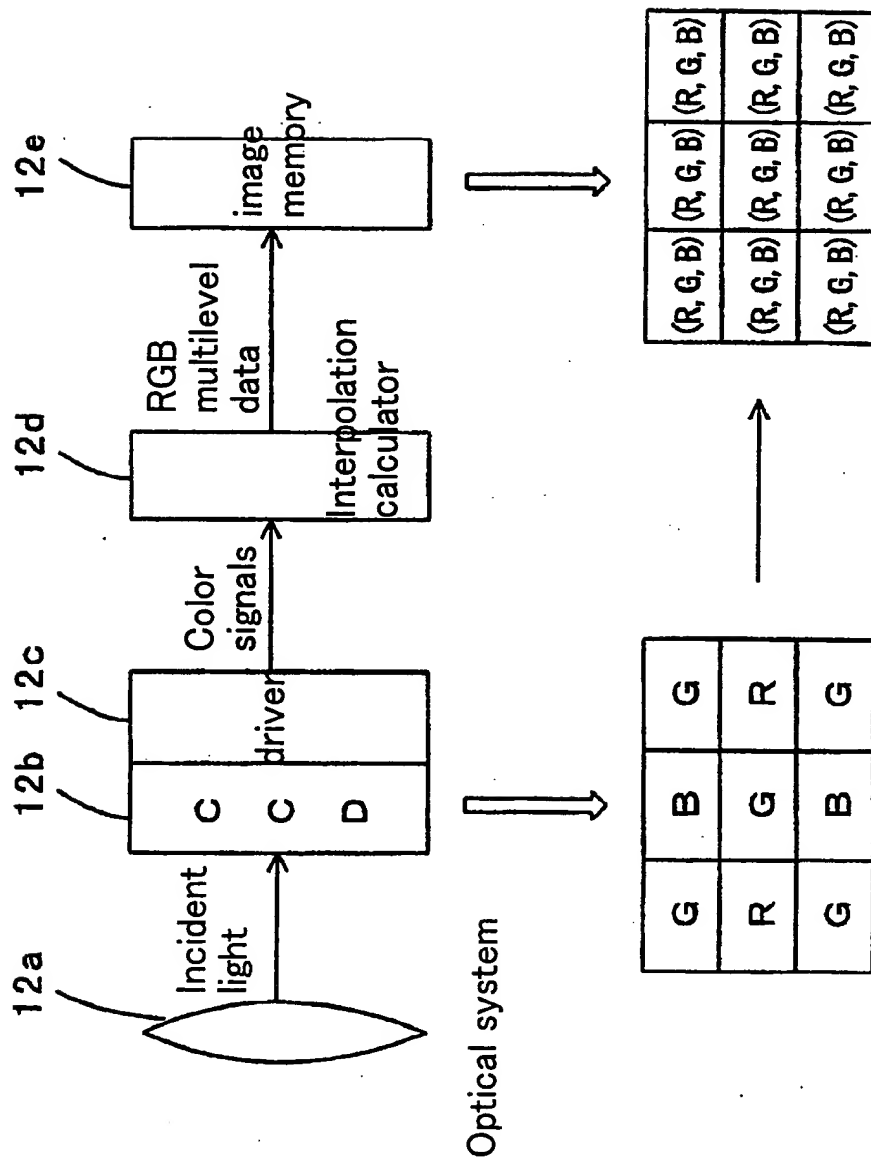


FIG.4

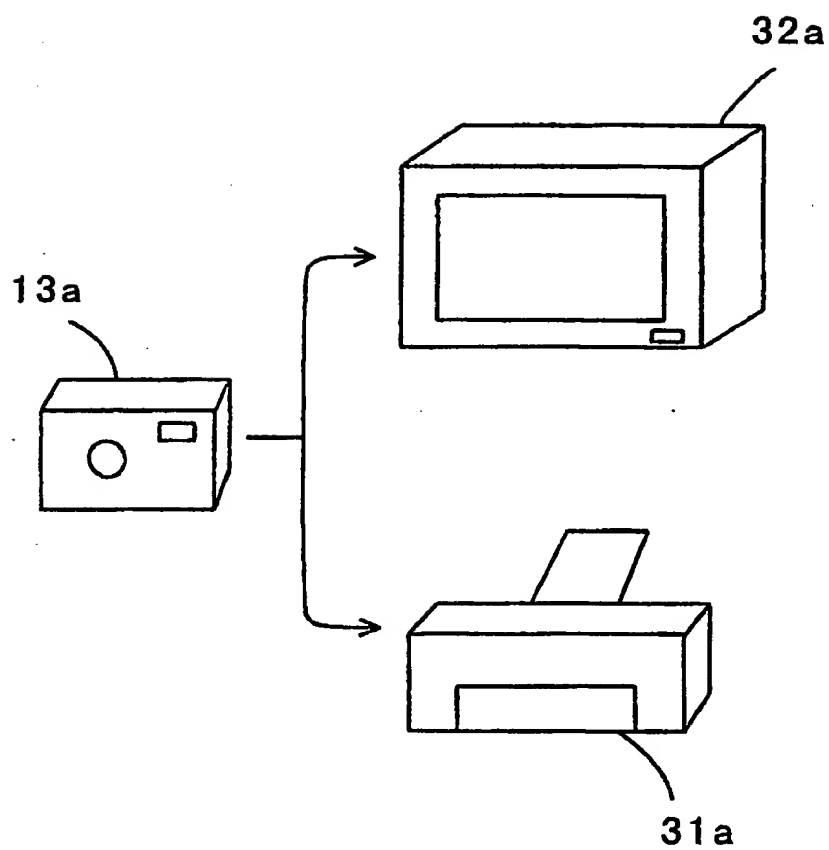
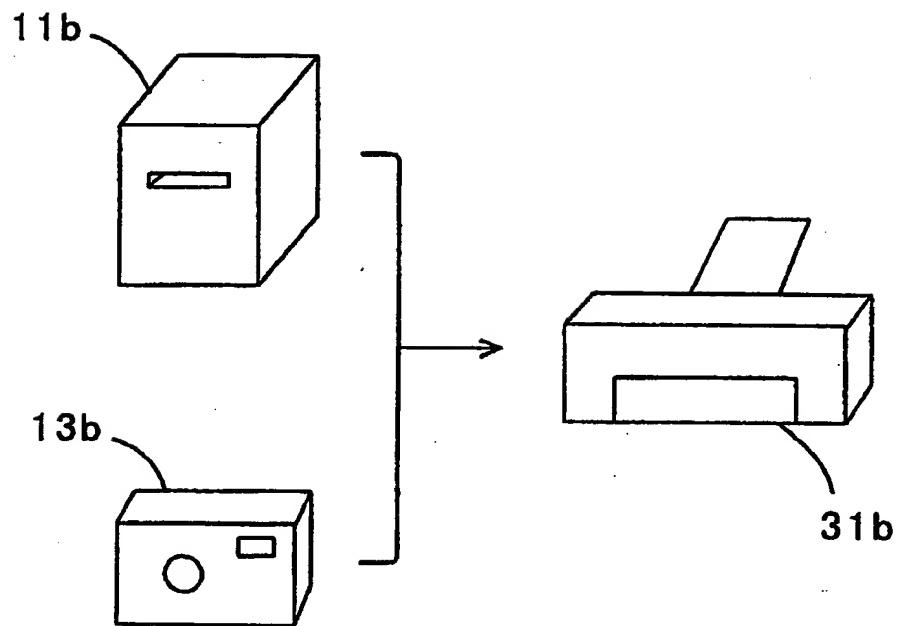
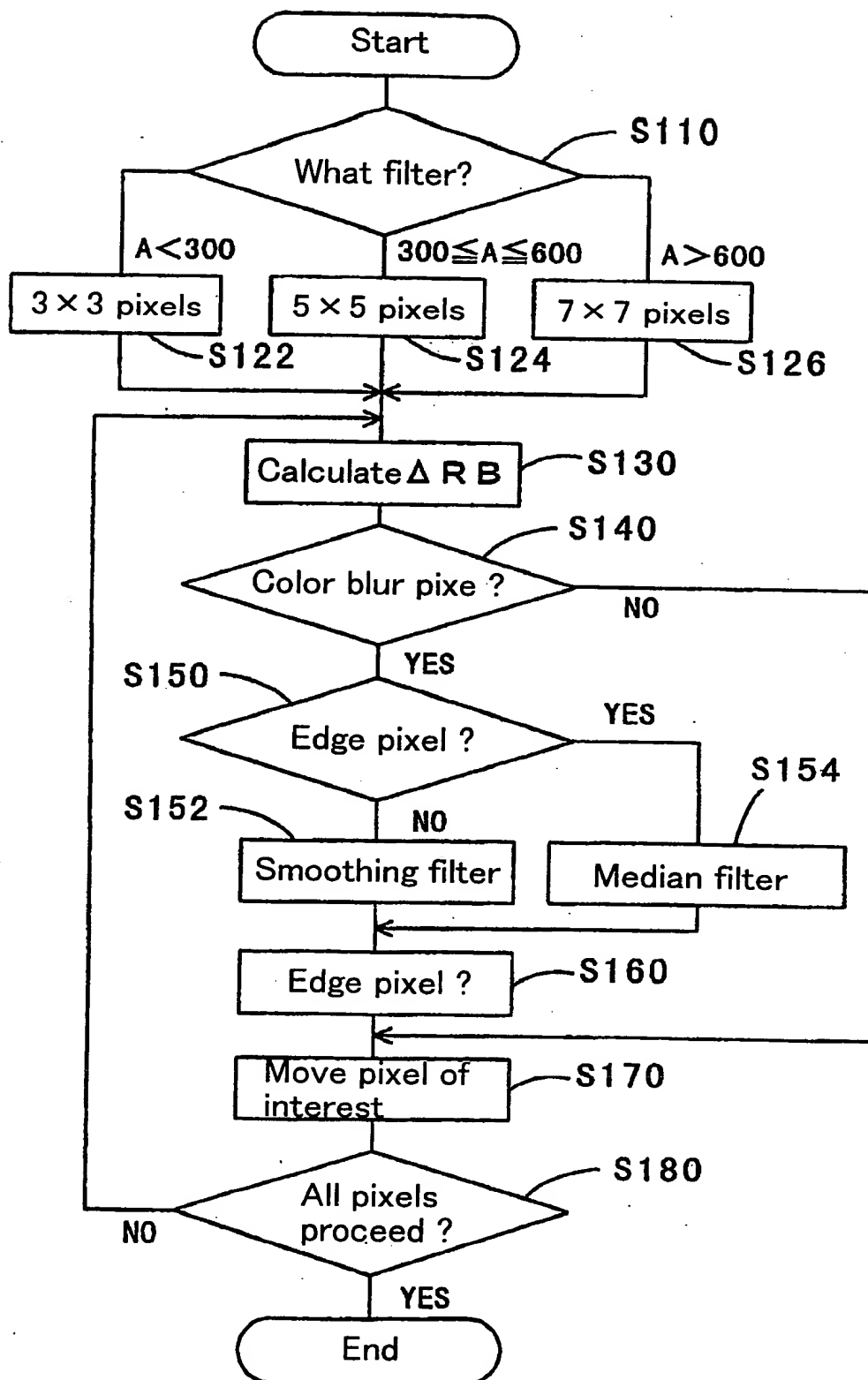


FIG.5



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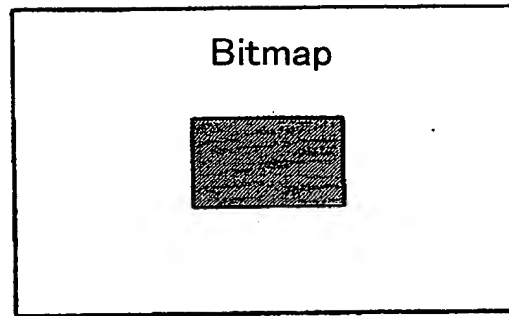
FIG.6



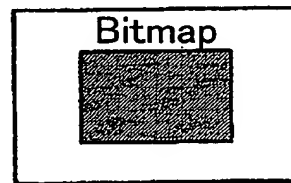
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FIG.7

(a)



(b)



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SECRET

FIG.8

(a)

1	1	1
1	1	1
1	1	1

3 × 3 pixels

(b)

1	1	1	1	1
1	1	1	1	1
1	1	1	1	1
1	1	1	1	1
1	1	1	1	1

5 × 5 pixels

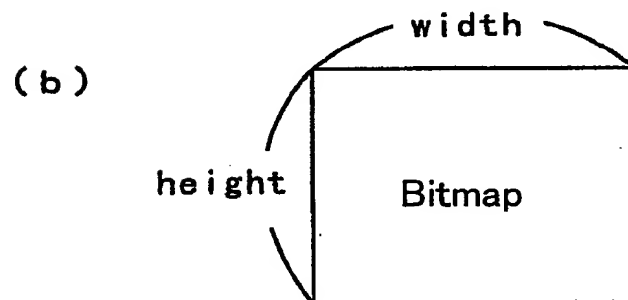
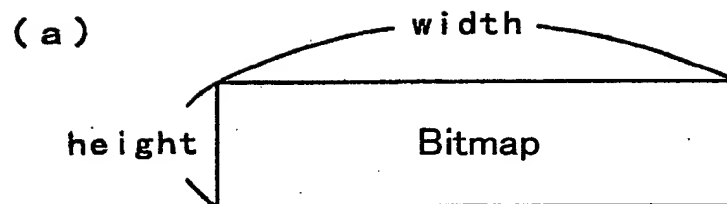
(c)

1	1	1	1	1	1	1
1	1	1	1	1	1	1
1	1	1	1	1	1	1
1	1	1	1	1	1	1
1	1	1	1	1	1	1
1	1	1	1	1	1	1
1	1	1	1	1	1	1

7 × 7 pixels

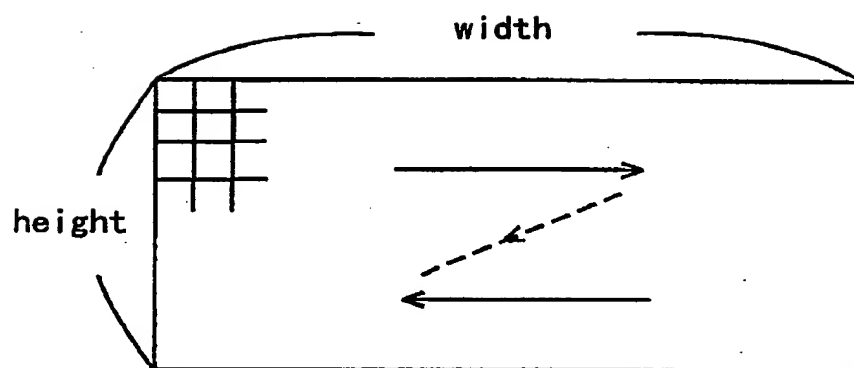
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FIG.9



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FIG.10



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$\Delta RB(x-1, y-1)$		$\Delta RB(x+1, y-1)$
	$\Delta RB(x, y)$	

[illegible]

FIG.12

$C1(x-1, y-1)$ $C2(x-1, y-1)$		
	$C1(x, y)$ $C2(x, y)$	

Patented

FIG.13

$C1(x-1, y-1)$ $C2(x-1, y-1)$	$C1(x, y-1)$ $C2(x, y-1)$	$C1(x+1, y-1)$ $C2(x+1, y-1)$
$C1(x-1, y)$ $C2(x-1, y)$	$C1(x, y)$ $C2(x, y)$	$C1(x+1, y)$ $C2(x+1, y)$
$C1(x-1, y+1)$ $C2(x-1, y+1)$	$C1(x, y+1)$ $C2(x, y+1)$	$C1(x+1, y+1)$ $C2(x+1, y+1)$

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Figure 1. The 1000 most abundant genes in the *Salmonella* genome. The genes are arranged in descending order of abundance. The x-axis represents the number of genes, and the y-axis represents the number of genes per 1000 genes. The data is presented as a bar chart with error bars.

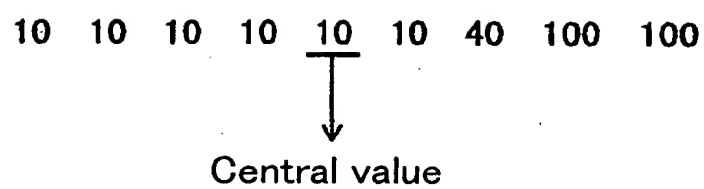
	$C1(x, y-1)$ $C2(x, y-1)$	
$C1(x-1, y)$ $C2(x-1, y)$	$C1(x, y)$ $C2(x, y)$	$C1(x+1, y)$ $C2(x+1, y)$
	$C1(x, y+1)$ $C2(x, y+1)$	

FIG.15

$C1(x-1, y-1)$ $C2(x-1, y-1)$		$C1(x+1, y-1)$ $C2(x+1, y-1)$
	$C1(x, y)$ $C2(x, y)$	
$C1(x-1, y+1)$ $C2(x-1, y+1)$		$C1(x+1, y+1)$ $C2(x+1, y+1)$

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FIG.17



Central value

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1	8	1
8	100	8
1	8	1

(total value of weighting coefficients...136)

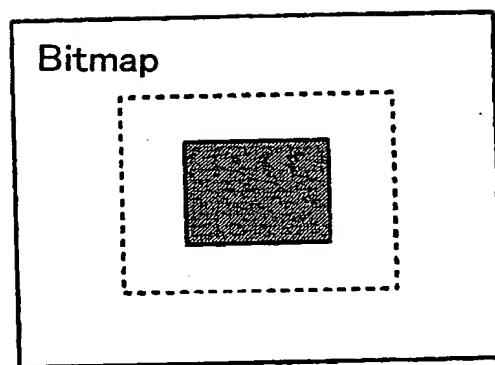
0	2	4	2	0
2	21	46	21	2
4	46	100	46	4
2	21	46	21	2
0	2	4	2	0

(total value of weighting coefficients...400)

0	1	3	4	3	1	0
1	6	18	25	18	6	1
3	18	50	71	50	18	3
4	25	71	100	71	25	4
3	18	50	71	50	18	3
1	6	18	25	18	6	1
0	1	3	4	3	1	0

(total value of weighting coefficients...900)

FIG.19



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FIG.20

FIG.21

FIG.22

FIG.23

